\_\_\_\_\_\_

Sequence Listing was accepted.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=7; day=21; hr=14; min=3; sec=21; ms=331; ]

# Validated By CRFValidator v 1.0.3

Application No: 10589054 Version No: 2.0

Input Set:

# Output Set:

**Started:** 2008-07-21 10:51:59.329

Finished: 2008-07-21 10:52:04.658

**Elapsed:** 0 hr(s) 0 min(s) 5 sec(s) 329 ms

Total Warnings: 136

Total Errors: 217

No. of SeqIDs Defined: 138

Actual SeqID Count: 138

| Error code |     | Error Description                                     |
|------------|-----|---|
| W 4        | 102 | Undefined organism found in <213> in SEQ ID (2)       |
| W 4        | 102 | Undefined organism found in <213> in SEQ ID (3)       |
| W 2        | 213 | Artificial or Unknown found in <213> in SEQ ID (5)    |
| E 2        | 257 | Invalid sequence data feature in <221> in SEQ ID (5)  |
| E 2        | 257 | Invalid sequence data feature in <221> in SEQ ID (5)  |
| E 2        | 257 | Invalid sequence data feature in <221> in SEQ ID (5)  |
| E 2        | 257 | Invalid sequence data feature in <221> in SEQ ID (5)  |
| E 2        | 257 | Invalid sequence data feature in <221> in SEQ ID (5)  |
| E 2        | 257 | Invalid sequence data feature in <221> in SEQ ID (5)  |
| W 2        | 213 | Artificial or Unknown found in <213> in SEQ ID (6)    |
| W 2        | 213 | Artificial or Unknown found in <213> in SEQ ID (7)    |
| W 2        | 213 | Artificial or Unknown found in <213> in SEQ ID (8)    |
| W 2        | 213 | Artificial or Unknown found in <213> in SEQ ID (9)    |
| W 2        | 213 | Artificial or Unknown found in <213> in SEQ ID (10)   |
| W 2        | 213 | Artificial or Unknown found in <213> in SEQ ID (11)   |
| W 2        | 213 | Artificial or Unknown found in <213> in SEQ ID (12)   |
| E 2        | 257 | Invalid sequence data feature in <221> in SEQ ID (12) |
| W 2        | 213 | Artificial or Unknown found in <213> in SEQ ID (13)   |
| E 2        | 257 | Invalid sequence data feature in <221> in SEQ ID (13) |
| W 2        | 213 | Artificial or Unknown found in <213> in SEQ ID (14)   |

### Input Set:

# Output Set:

**Started:** 2008-07-21 10:51:59.329

Finished: 2008-07-21 10:52:04.658

**Elapsed:** 0 hr(s) 0 min(s) 5 sec(s) 329 ms

Total Warnings: 136

Total Errors: 217

No. of SeqIDs Defined: 138

Actual SeqID Count: 138

| Error code |     | Error Description  |
|------------|-----|--|
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (15)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (16)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (17)  |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (17)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (18)  |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (18)  |
| M          | 213 | Artificial or Unknown found in <213> in SEQ ID (19)  |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (19)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (20)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (21)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (22)  |
| M          | 213 | Artificial or Unknown found in <213> in SEQ ID (23)  |
| W          | 213 | Artificial or Unknown found in <213> in SEQ ID (24) This error has occured more than 20 times, will not be displayed |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (29)  |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (29)  |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (29)  |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (29)  |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (29)  |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (29)  |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (29)  |
| E          | 257 | Invalid sequence data feature in <221> in SEQ ID (29)  |
| Ε          | 257 | Invalid sequence data feature in <221> in SEQ ID (29)  |

### Input Set:

# Output Set:

**Started:** 2008-07-21 10:51:59.329

Finished: 2008-07-21 10:52:04.658

**Elapsed:** 0 hr(s) 0 min(s) 5 sec(s) 329 ms

Total Warnings: 136

Total Errors: 217

No. of SeqIDs Defined: 138

Actual SeqID Count: 138

Error code Error Description

### SEQUENCE LISTING

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<110> SOARES, CHRISTOPHER J.
      HANLEY, MICHAEL R.
      LEWIS, DIANA YVONNE
      PARKES, DAVID GEOFFREY
      JODKA, CAROLYN MARIE
      PRICKETT, KATHRYN S.
      GHOSH, SOUMITRA
      MACK, CHRISTINE MARIE
      LIN, QING
<120> AMYLIN FAMILY PEPTIDES AND METHODS FOR MAKING AND USING
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<130> 0105US-UTL2
<140> 10589054
<141> 2008-07-21
<150> PCT/US05/004631
<151> 2005-02-11
<150> 60/550,447
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<213> Oncorhynchus sp.

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<211> 32

<212> PRT

<213> Homo sapiens

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<212> PRT

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 specification as filed for detailed
 description of substitutions and preferred
 embodiments

<220>

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1 5

1

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<222> (12)
<223> Glu or Phe
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<223> His or Aib
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                                                          15
 1
                                      10
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             20
                                  25
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Lys (PEG5000)
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<222> (16)
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<222> (17)
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<223> Thr or Asn
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<223> Tyr, Val, Phe, Leu, or Met
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<222> (25)..(28)
<223> This region may encompass 1-4 Any amino acid
      residues
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                                     10
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             20
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      C-terminal amino acid sequence
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<221> MOD\_RES

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20
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<221> MOD_RES
<222> (9)
<223> Leu or Thr
<220>
<221> MOD_RES
<222> (10)
<223> Gly, His, or
```